

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/563,310  
Source: IFWP  
Date Processed by STIC: 2/21/07

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1). INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>) , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):  
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

## Raw Sequence Listing Error Summary

### ERROR DETECTED

### SUGGESTED CORRECTION

### SERIAL NUMBER:

10/563,310

**ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE**

- |    |  |  |
|----|--|--|
| 1  | ____ Wrapped Nucleics<br>____ Wrapped Aminos | The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."   |
| 2  | ____ Invalid Line Length                     | The rules require that a line <b>not exceed</b> 72 characters in length. This includes white spaces.   |
| 3  | ____ Misaligned Amino Numbering              | The numbering under each 5 <sup>th</sup> amino acid is misaligned. Do <b>not</b> use tab codes between numbers; use <b>space characters</b> , instead.   |
| 4  | ____ Non-ASCII                               | The submitted file was <b>not saved</b> in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is <b>saved in ASCII text</b> .  |
| 5  | ____ Variable Length                         | Sequence(s) ____ contain n's or Xaa's representing more than one residue. <b>Per Sequence Rules, each n or Xaa can only represent a single residue.</b> Please present the <b>maximum</b> number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.  |
| 6  | ____ PatentIn 2.0 "bug"                      | A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. <b>This applies to the mandatory &lt;220&gt;-&lt;223&gt; sections for Artificial or Unknown sequences.</b>   |
| 7  | ____ Skipped Sequences (OLD RULES)           | Sequence(s) ____ missing. If intentional, please insert the following lines for <b>each</b> skipped sequence:<br>(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)<br>(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)<br>(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)<br>This sequence is intentionally skipped<br>Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to <b>include the skipped sequences</b> . |
| 8  | ____ Skipped Sequences (NEW RULES)           | Sequence(s) ____ missing. If intentional, please insert the following lines for <b>each</b> skipped sequence.<br><210> sequence id number<br><400> sequence id number<br>000   |
| 9  | ____ Use of n's or Xaa's (NEW RULES)         | Use of n's and/or Xaa's have been detected in the Sequence Listing.<br>Per 1.823 of Sequence Rules, use of <220>-<223> is <b>MANDATORY</b> if n's or Xaa's are present.<br>In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.  |
| 10 | ____ Invalid <213> Response                  | Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is <b>required</b> when <213> response is Unknown or is Artificial Sequence. (see item 11 below)  |
| 11 | ____ Use of <220>                            | Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is <b>MANDATORY</b> if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules   |
| 12 | ____ PatentIn 2.0 "bug"                      | Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.  |
| 13 | ____ Misuse of n/Xaa                         | "n" <b>can only represent a single nucleotide</b> ; "Xaa" <b>can only represent a single amino acid</b>  |



IFWO

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/563,310

DATE: 02/21/2007  
TIME: 14:14:40

Input Set : A:\21013981.ST25.txt  
Output Set: N:\CRF4\02212007\J563310.raw

3 <110> APPLICANT: Affibody AB  
 5 <120> TITLE OF INVENTION: Polypeptides having binding affinity for HER2  
 7 <130> FILE REFERENCE: P106236PCT  
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/563,310  
 C--> 10 <141> CURRENT FILING DATE: 2005-12-29  
 12 <150> PRIOR APPLICATION NUMBER: SE0301987-4  
 13 <151> PRIOR FILING DATE: 2003-07-04  
 15 <150> PRIOR APPLICATION NUMBER: SE0400275-4  
 16 <151> PRIOR FILING DATE: 2004-02-09  
 18 <160> NUMBER OF SEQ ID NOS: 79  
 20 <170> SOFTWARE: PatentIn version 3.3  
 22 <210> SEQ ID NO: 1  
 23 <211> LENGTH: 58  
 24 <212> TYPE: PRT  
 25 <213> ORGANISM: Synthetic polypeptide  
 27 <400> SEQUENCE: 1  
 29 Val Asp Asn Lys Phe Asn Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile  
 30 1 5 .10 15  
 33 Leu His Leu Pro Asn Leu Asn Glu Glu Gln Arg Asn Ala Phe Ile Gln  
 34 20 25 30  
 37 Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
 38 35 40 45  
 41 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
 42 50 55  
 45 <210> SEQ ID NO: 2  
 46 <211> LENGTH: 58  
 47 <212> TYPE: PRT  
 48 <213> ORGANISM: Synthetic polypeptide  
 50 <400> SEQUENCE: 2  
 52 Val Asp Asn Lys Phe Asn Lys Glu Leu Arg Gln Ala Tyr Trp Glu Ile  
 53 1 5 10 15  
 56 Gln Ala Leu Pro Asn Leu Asn Trp Thr Gln Ser Arg Ala Phe Ile Arg  
 57 20 25 30  
 60 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
 61 35 40 45  
 64 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
 65 50 55  
 68 <210> SEQ ID NO: 3  
 69 <211> LENGTH: 58  
 70 <212> TYPE: PRT  
 71 <213> ORGANISM: Synthetic polypeptide  
 73 <400> SEQUENCE: 3  
 75 Val Asp Asn Lys Phe Asn Lys Glu Pro Lys Thr Ala Tyr Trp Glu Ile

*See pg 1-5*  
*invalid response*  
*see item 10 on Error summary*  
*sheet*

*Does Not Comply*  
*Corrected Diskette Needed*

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/563,310

DATE: 02/21/2007  
TIME: 14:14:40

Input Set : A:\21013981.ST25.txt  
Output Set: N:\CRF4\02212007\J563310.raw

76 1 5 10 15  
79 Val Lys Leu Pro Asn Leu Asn Pro Glu Gln Arg Arg Ala Phe Ile Arg  
80 20 25 30  
83 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
84 35 40 45  
87 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
88 50 55  
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92 <211> LENGTH: 58  
93 <212> TYPE: PRT  
94 <213> ORGANISM: Synthetic polypeptide  
96 <400> SEQUENCE: 4  
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99 1 5 10 15  
102 Gln Arg Leu Pro Asn Leu Asn Asn Lys Gln Lys Ala Ala Phe Ile Arg  
103 20 25 30  
106 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
107 35 40 45  
110 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
111 50 55  
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125 Tyr Asn Leu Pro Asn Leu Asn Arg Ala Gln Met Arg Ala Phe Ile Arg  
126 20 25 30  
129 Ser Leu Ser Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
130 35 40 45  
133 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
134 50 55  
137 <210> SEQ ID NO: 6  
138 <211> LENGTH: 58  
139 <212> TYPE: PRT  
140 <213> ORGANISM: SYNTHETIC POLYPEPTIDE  
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148 Val Lys Leu Pro Asn Leu Asn Pro Arg Gln Lys Arg Ala Phe Ile Arg  
149 20 25 30  
152 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
153 35 40 45  
156 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
157 50 55  
160 <210> SEQ ID NO: 7  
161 <211> LENGTH: 58  
162 <212> TYPE: PRT

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/563,310

DATE: 02/21/2007  
TIME: 14:14:40

Input Set : A:\21013981.ST25.txt  
Output Set: N:\CRF4\02212007\J563310.raw

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168 1 5 10 15  
171 Val Leu Leu Pro Asn Leu Asn Arg Arg Gln Ser Arg Ala Phe Ile Arg  
172 20 25 30  
175 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
176 35 40 45  
179 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
180 50 55  
183 <210> SEQ ID NO: 8  
184 <211> LENGTH: 58  
185 <212> TYPE: PRT  
186 <213> ORGANISM: **SYNTHETIC POLYPEPTIDE**  
188 <400> SEQUENCE: 8  
190 Val Asp Asn Lys Phe Asn Lys Glu Met Arg His Ala Tyr Trp Glu Ile  
191 1 5 10 15  
194 Ala Thr Leu Pro Asn Leu Asn Asn Val Gln Lys Arg Ala Phe Ile Arg  
195 20 25 30  
198 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
199 35 40 45  
202 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
203 50 55  
206 <210> SEQ ID NO: 9  
207 <211> LENGTH: 58  
208 <212> TYPE: PRT  
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214 1 5 10 15  
217 Val Leu Leu Pro Asn Leu Asn Pro Gly Gln Ile Arg Ala Phe Ile Arg  
218 20 25 30  
221 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
222 35 40 45  
225 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
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231 <212> TYPE: PRT  
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240 Val Leu Leu Pro Asn Leu Asn Thr Trp Gln Ile Arg Ala Phe Ile Arg  
241 20 25 30  
244 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
245 35 40 45  
248 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
249 50 55

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/563,310

DATE: 02/21/2007  
TIME: 14:14:40

Input Set : A:\21013981.ST25.txt  
Output Set: N:\CRF4\02212007\J563310.raw

252 <210> SEQ ID NO: 11  
253 <211> LENGTH: 58  
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260 1 5 10 15  
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264 20 25 30  
267 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
268 35 40 45  
271 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
272 50 55  
275 <210> SEQ ID NO: 12  
276 <211> LENGTH: 58  
277 <212> TYPE: PRT  
278 <213> ORGANISM: SYNTHETIC POLYPEPTIDE  
280 <400> SEQUENCE: 12  
282 Val Asp Asn Lys Phe Asn Lys Glu Met Arg Asn Ala Tyr Trp Glu Ile  
283 1 5 10 15  
286 Ala Leu Leu Pro Asn Leu Asn Asn Gln Gln Lys Arg Ala Phe Ile Arg  
287 20 25 30  
290 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
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300 <212> TYPE: PRT  
301 <213> ORGANISM: SYNTHETIC POLYPEPTIDE  
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306 1 5 10 15  
309 Val Gly Leu Pro Asn Leu Asn His Phe Gln Val Arg Ala Phe Ile Arg  
310 20 25 30  
313 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala  
314 35 40 45  
317 Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys  
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323 <212> TYPE: PRT  
324 <213> ORGANISM: SYNTHETIC POLYPEPTIDE  
326 <400> SEQUENCE: 14  
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332 Val Leu Leu Pro Asn Leu Asn Arg Trp Gln Ile Arg Ala Phe Ile Arg  
333 20 25 30  
336 Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/563,310

DATE: 02/21/2007  
TIME: 14:14:40

Input Set : A:\21013981.ST25.txt  
Output Set: N:\CRF4\02212007\J563310.raw

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355	Ala Leu Leu Pro Asn Leu Asn Asn Met Gln Lys Arg Ala Phe Ile Arg		
356	20 25 30		
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379	20 25 30		
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402	20 25 30		
405	Ser Leu Tyr Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala		
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421	1 5 10 15		

*Please correct this error  
in subsequent sequences.*

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/563,310

DATE: 02/21/2007

TIME: 14:14:41

Input Set : A:\21013981.ST25.txt

Output Set: N:\CRF4\02212007\J563310.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date